

Simulation of Shock Waves by Smoothed Particle Hydrodynamics

Mohsen Nejad-Asghar

School of Physics, Damghan University of Basic Sciences, Damghan, Iran

nasghar@dubs.ac.ir

ABSTRACT

Isothermal and adiabatic shocks, which are produced from fast expansion of the gas, is simulated with smoothed particle hydrodynamics (SPH). The results are compared with the analytic solutions. The algorithm of the program is explained and the package, which is written in Fortran, is presented in the appendix of this paper. It is possible to change (to complete) the program for a wide variety of applications ranging from astrophysics to fluid mechanics.

Subject headings: Hydrodynamics, methods: numerical, ISM: evolution

1. Introduction

Gas dynamical processes are believed to play an important role in the evolution of astrophysical systems on all length scales. Smoothed particle hydrodynamics (SPH) is a powerful gridless particle method to solve complex fluid-dynamical problems. SPH has a number of attractive features such as its low numerical diffusion in comparison to grid based methods. An adequate scenario for SPH application is the unbound astrophysical problems, especially on the shock propagation (see, e.g., Liu & Liu 2003). In this way, the basic principles of the SPH is written in this paper and the simulation of isothermal and adiabatic shocks are applied to test the ability of this numerical simulation to produce known analytic solutions.

The program is written in Fortran and is highly portable. This package supports only calculations for isothermal and adiabatic shock waves. It is possible to change (to complete) the program for a wide variety of applications ranging from astrophysics to fluid mechanics. The program is written in modular form, in the hope that it will provide a useful tool. I ask only that:

- If you publish results obtained using some parts of this program, please consider acknowledging the source of the package.

- If you discover any errors in the program or documentation, please promptly communicate the to the author.

2. Formulation of Shock Waves

An extremely important problem is the behavior of gases subjected to compression waves. This happens very often in the cases of astrophysical interests. For example, a small region of gas suddenly heated by the liberation of energy will expand into its surroundings. The surroundings will be pushed and compressed. Conservation of mass, momentum, and energy across a shock front is given by the Rankine-Hugoniot conditions (Dyson & Williams 1997)

$$\rho_1 v_1 = \rho_2 v_2 \quad (1)$$

$$\rho_1 v_1^2 + K \rho_1^\gamma = \rho_2 v_2^2 + K \rho_2^\gamma \quad (2)$$

$$\frac{1}{2} v_1^2 + \frac{\gamma}{\gamma-1} K \rho_1^{\gamma-1} = \frac{1}{2} v_2^2 + \frac{\gamma}{\gamma-1} K \rho_2^{\gamma-1} + Q \quad (3)$$

where the equation of state, $p = K \rho^\gamma$, is used. In adiabatic case, we have $Q = 0$, and for isothermal shocks, we will set $\gamma = 1$.

We would interested to consider the collision of two gas sheets with velocities v_0 in the rest frame of the laboratory. In this reference frame, the post-shock will be at rest and the pre-shock velocity is given by $v_1 = v_0 + v_2$, where v_2 is the shock front velocity. Combining equations (1)-(3), we have

$$v_2 = a_0 \left[-\frac{b}{2} + \sqrt{1 + \frac{b^2}{4} + (\gamma-1) \left(\frac{M_0^2}{2} - q \right)} \right] \quad (4)$$

where $a_0 \equiv \gamma K \rho_1^{\gamma-1}$ is the sound speed, $M_0 \equiv v_0/a_0$ is the Mach number, b and q are defined as

$$b \equiv \frac{3-\gamma}{2} M_0 + \frac{\gamma-1}{M_0} q \quad ; \quad q \equiv \frac{Q}{a_0^2}. \quad (5)$$

Substituting (4) into equation (1), density of the post-shock is given by

$$\rho_2 = \rho_1 \left\{ 1 + \frac{M_0}{\left[-\frac{b}{2} + \sqrt{1 + \frac{b^2}{4} + (\gamma-1) \left(\frac{M_0^2}{2} - q \right)} \right]} \right\}. \quad (6)$$

3. SPH Equations

The smoothed particle hydrodynamics was invented to simulate nonaxisymmetric phenomena in astrophysics (Lucy 1977, Gingold & Monaghan 1977). In this method, fluid is represented by N discrete but extended/smoothed particles (i.e. Lagrangian sample points). The particles are overlapping, so that all the physical quantities involved can be treated as continuous functions both in space and time. Overlapping is represented by the kernel function, $W_{ab} \equiv W(\mathbf{r}_a - \mathbf{r}_b, h_{ab})$, where $h_{ab} \equiv (h_a + h_b)/2$ is the mean smoothing length of two particles a and b . The continuity, momentum and energy equation of particle a are (Monaghan 1992)

$$\rho_a = \sum_b m_b W_{ab} \quad (7)$$

$$\frac{d\mathbf{v}_a}{dt} = - \sum_b m_b \left(\frac{p_a}{\rho_a} + \frac{p_b}{\rho_b} + \Pi_{ab} \right) \nabla_a W_{ab} \quad (8)$$

$$\frac{du_a}{dt} = \frac{1}{2} \sum_b m_b \left(\frac{p_a}{\rho_a} + \frac{p_b}{\rho_b} + \Pi_{ab} \right) \mathbf{v}_{ab} \cdot \nabla_a W_{ab} \quad (9)$$

where $\mathbf{v}_{ab} \equiv \mathbf{v}_a - \mathbf{v}_b$ and

$$\Pi_{ab} = \begin{cases} \frac{\alpha v_{sig} \mu_{ab} + \beta \mu_{ab}^2}{\bar{\rho}_{ab}}, & \text{if } \mathbf{v}_{ab} \cdot \mathbf{r}_{ab} < 0, \\ 0, & \text{otherwise,} \end{cases} \quad (10)$$

is the artificial viscosity between particles a and b , where $\bar{\rho}_{ab} = \frac{1}{2}(\rho_a + \rho_b)$ is an average density, $\alpha \sim 1$ and $\beta \sim 2$ are the artificial coefficients, and μ_{ab} is defined as its usual form

$$\mu_{ab} = - \frac{\mathbf{v}_{ab} \cdot \mathbf{r}_{ab}}{\bar{h}_{ab}} \frac{1}{r_{ab}^2 / \bar{h}_{ab}^2 + \eta^2} \quad (11)$$

with $\eta \sim 0.1$ and $\bar{h}_{ab} = \frac{1}{2}(h_a + h_b)$. The signal velocity, v_{sig} , is

$$v_{sig} = \frac{1}{2}(c_a + c_b) \quad (12)$$

where c_a and c_b are the sound speed of particles. The SPH equations are integrated using the smallest time-steps

$$\Delta t_a = C_{cour} \text{MIN} \left[\frac{h_a}{|\mathbf{v}_a|}, \left(\frac{h_1}{|\mathbf{a}_1|} \right)^{0.5}, \frac{u_a}{|du_a/dt|}, \frac{h_a}{|dh_a/dt|}, \frac{\rho_a}{|d\rho_a/dt|} \right] \quad (13)$$

where $C_{cour} \sim 0.25$ is the Courant number.

4. Results and Prospects

The chosen physical scales for length and time are $[l] = 3.0 \times 10^{16} m$ and $[t] = 3.0 \times 10^{13} s$, respectively, so the velocity unit is approximately $1 km.s^{-1}$. The gravity constant is set $G = 1$ for which the calculated mass unit is $[m] = 4.5 \times 10^{32} kg$. There is considered two equal one dimensional sheets with extension $x = 0.1[l]$, which have initial uniform density and temperature of $\sim 4.5 \times 10^8 m^{-3}$ and $\sim 10 K$, respectively.

Particles with a positive x-coordinate are given an initial negative velocity of Mach 5, and those with a negative x-coordinate are given a Mach 5 velocity in the opposite direction. In adiabatic shock, with $M_0 = 5$, the post-shock density must be 2.9, which is obtained from analytic solution (6) with $Q = 0$ and $\gamma = 2$. The Results of adiabatic shock are shown in Fig. 1-3. In isothermal shock, with $M_0 = 5$, the post-shock density must be 26.9, which is obtained from analytic solution Equ. (6) with $\gamma = 1$. The Results of isothermal shock are shown in Fig. 4-5. Algorithm of the program is shown in Fig. 6.

REFERENCES

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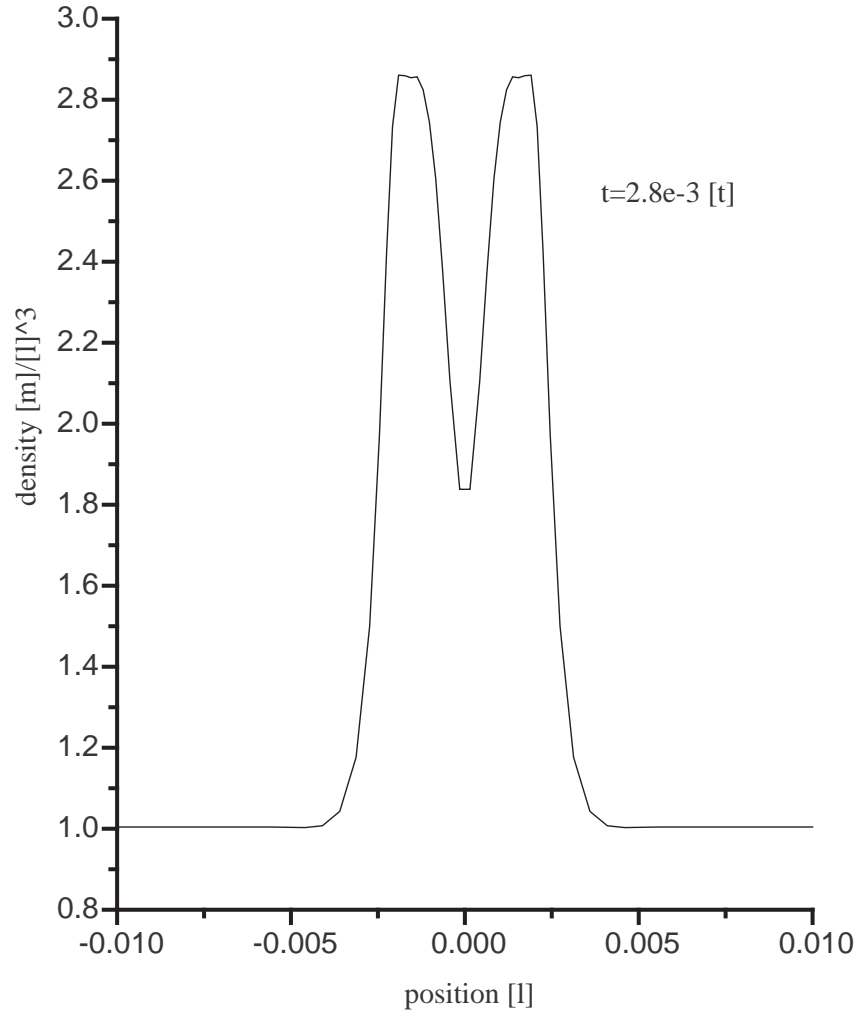


Fig. 1.— The density of adiabatic shock, with $M_0 = 5$, $Q = 0$, and $\gamma = 2$.

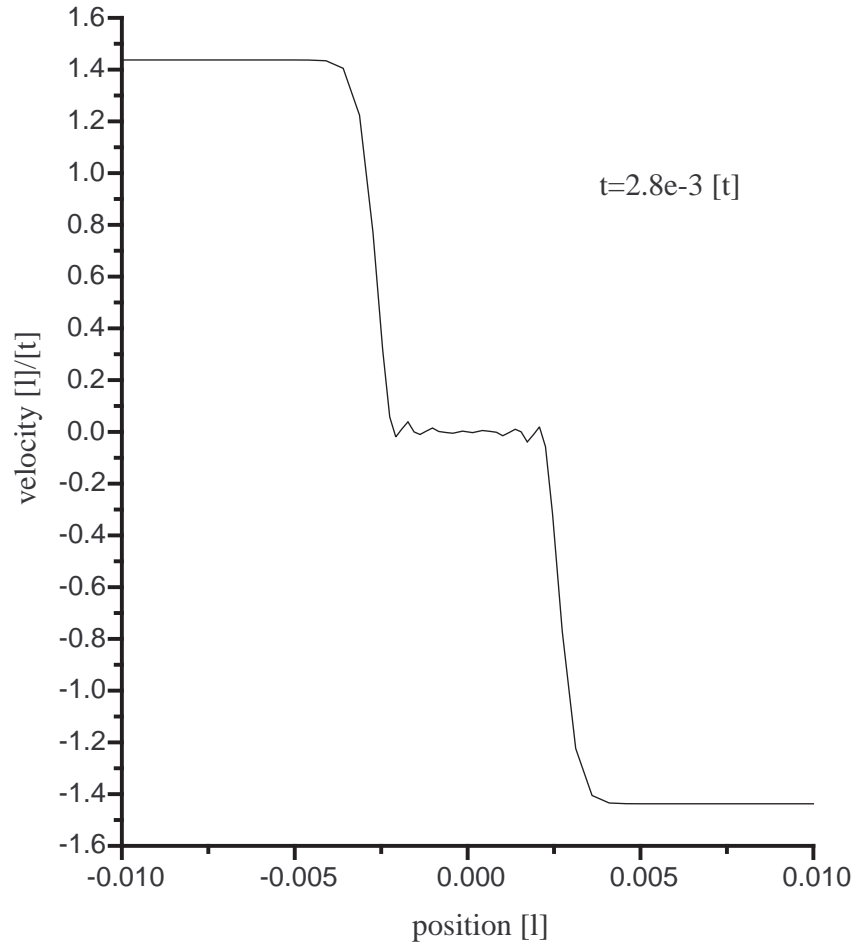


Fig. 2.— The velocity of adiabatic shock, with $M_0 = 5$, $Q = 0$, and $\gamma = 2$.

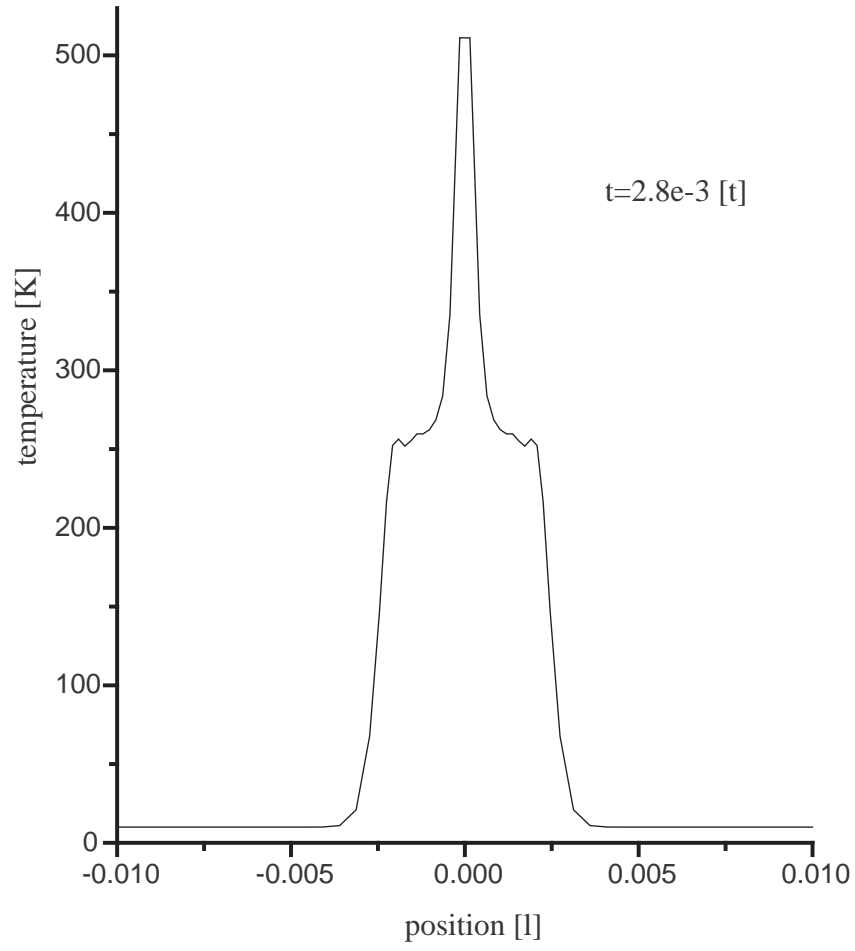


Fig. 3.— The temperature of adiabatic shock, with $M_0 = 5$, $Q = 0$, and $\gamma = 2$.

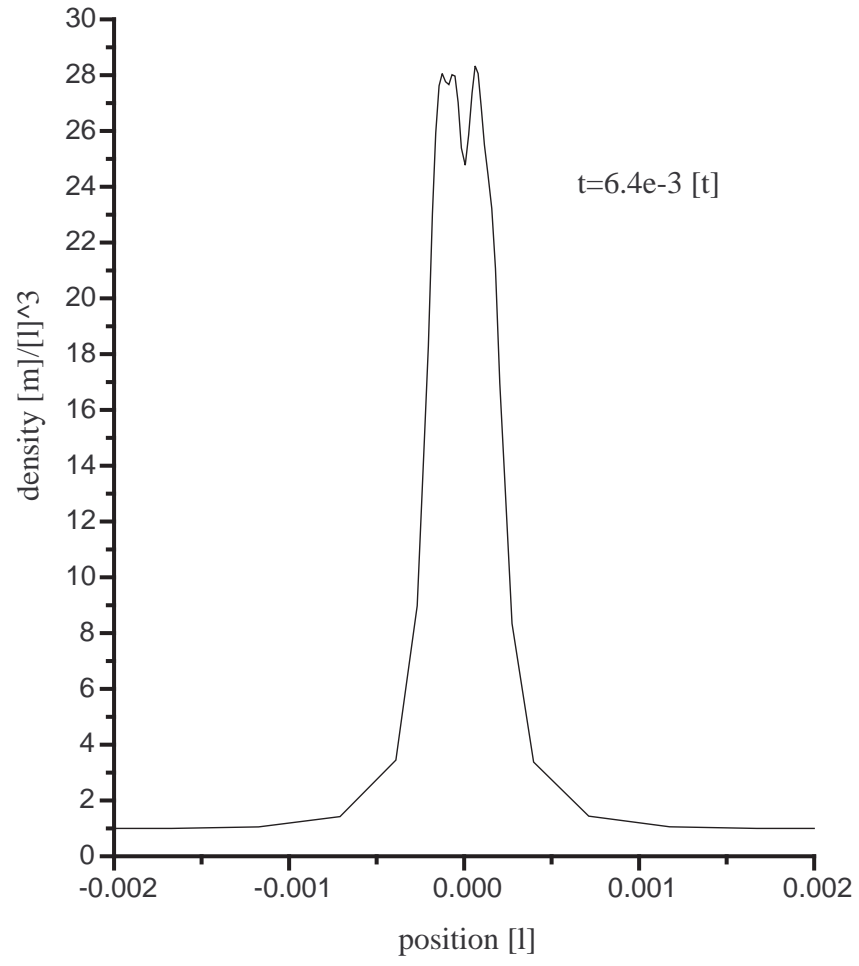


Fig. 4.— The density of isothermal shock, with $M_0 = 5$ and $\gamma = 1$.

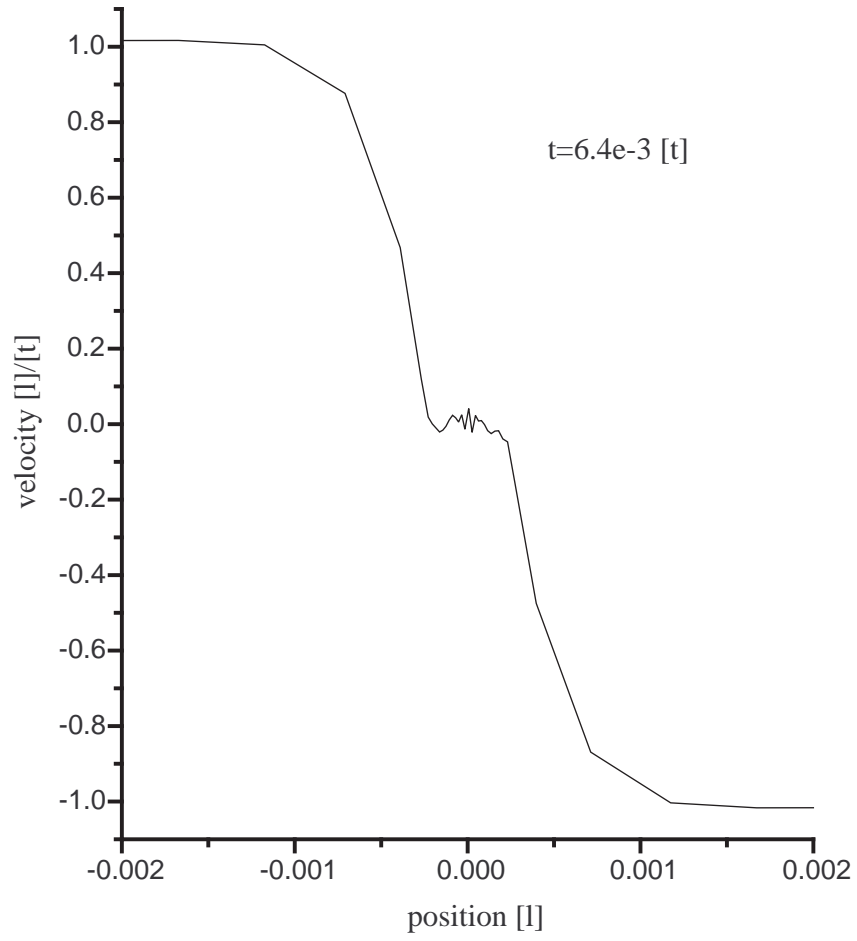


Fig. 5.— The velocity of isothermal shock, with $M_0 = 5$ and $\gamma = 1$.

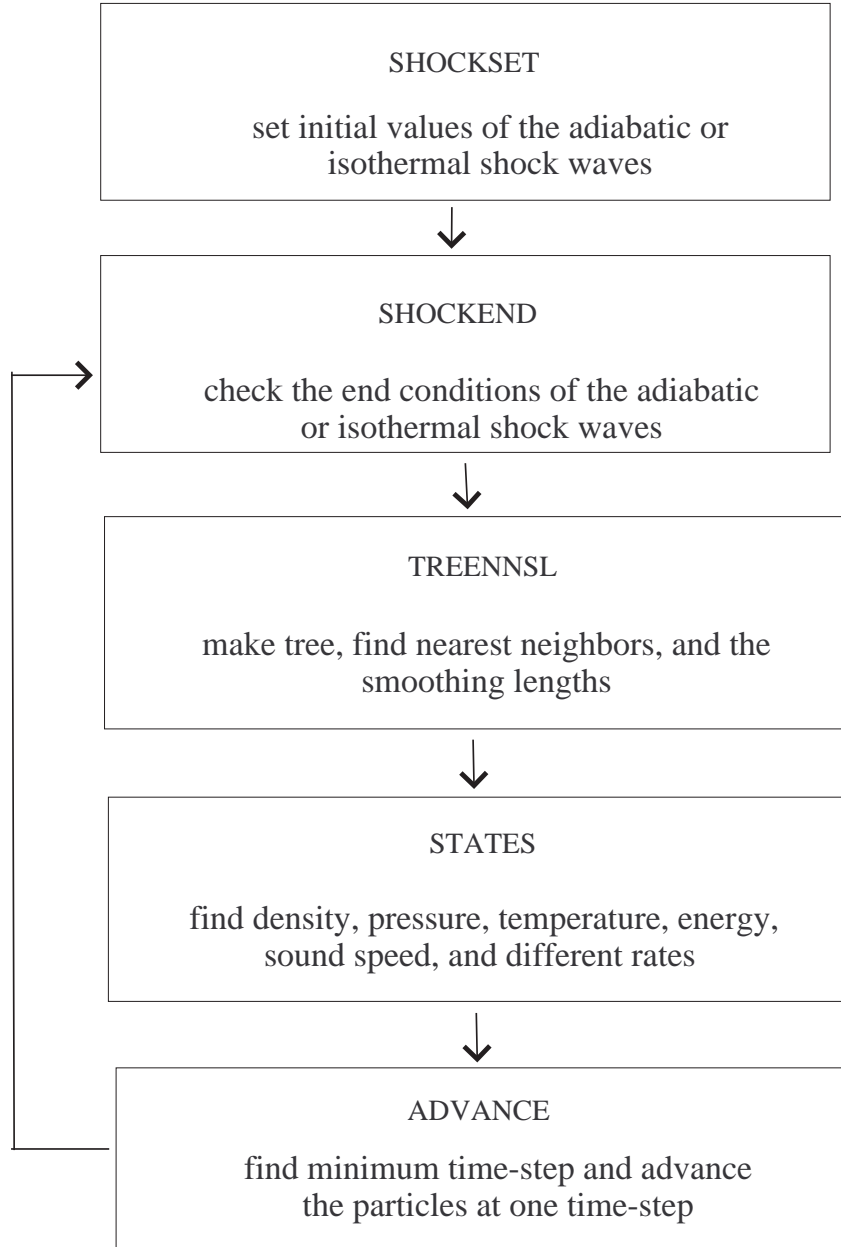


Fig. 6.— Algorithm of the smoothed particle hydrodynamics for simulation of isothermal and adiabatic shocks.

```
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!      This program is provided to simulate the adiabatic
!      and isothermal shock waves
!      *****
!      Mohsen Nejad-Asghar
!      nasghar@dubs.ac.ir
!      January 2006
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      PROGRAM SHOCK
      INCLUDE 'param.inc'
      PRINT*, 'isothermal or adiabatic shock?'
      PRINT*, 'adiabatic=1'
      PRINT*, 'isothermal=2'
      READ(*,*) isorad
      CALL SHOCKSET
      CALL SCENARIOS
      ! investigate the end condition of simulation
10    CALL SHOCKEND
      ! advance system at one time-step
      CALL ADVANCE
      GOTO 10
      END PROGRAM SHOCK

!=====
!////////////////////
!=====
      SUBROUTINE SHOCKSET
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!      This subroutine generates initial particle information for
!      adiabatic or isothermal one dimensional shock
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      INCLUDE 'param.inc'
      REAL extx, delx, mach
      nbody=400
      IF(nbody > maxn)
&      CALL ERROR('SHOCKSET: SPH number is very large')
      mxcell=nsubc*nbody
      node=nbody+mxcell
      incell=nbody+1
      ! units of length, time, and mass
      ul=3.0e16
      ut=3.0e13
      um=4.5e32
      ! extension of each sheet in x direction (ul)
      extx=0.1
      ! positions of SPH particles
      delx=2.0*extx/nbody
      DO i=1, nbody
        pos(i,1)=extx-i*delx+delx/2.0
      END DO
      ! density of SPH particles (um/ul^3)
      den=1.0
      ! temperature of SPH particles (K)
      temp=10.0
      ! masses of SPH particles (um)
      DO i=1, nbody
        mass(i)=delx*den(i)
      END DO
      ! molecular weight relative to the mass of hydrogen
      xmu=2.0
      ! hydrogen mass
      xmh=1.67e-27
      ! Boltzman constant
      xkb=1.38e-23
      xKK=(xkb/(xmu*xmh))/(ul/ut)**2
      IF(isorad == 1)THEN
        ! polytropic index (adiabatic case)
        gamma=2.0
        ! energy of SPH particles
        u=xKK*temp/(gamma-1)
      ELSEIF(isorad == 2)THEN
        ! polytropic index (isothermal case)
        gamma=1.0
      ENDIF
      ! sound speed
      sound=SQRT(gamma*xKK*temp)
      ! Mach number
      mach=5.0
      IF(isorad == 1)THEN
        ! relative density after simulation for adiabatic case
        cons=SQRT(16.0+((3-gamma)**2+8*(gamma-1))*mach*mach)
```

[illegible]

```

INCLUDE 'param.inc'
! skf--> smoothing kernel function?
!      =1: Gauss kernel (Gingold & Monaghan 1981)
!      =2: spline-base kernel (Monaghan 1985)
!      =3: Quintic kernel (Morris 1997)
skf=2
! nnssl--> nearest neighbors and smoothing length?
!      =1: fixed smoothing length
!      =2: variable smoothing length
nnssl=1
! dsm--> density summation method?
!      =1: summation model without continuity
!      =2: use continuity equation
dsm=1
! the artificial shear viscosity?
alphas=1.0
! the artificial bulk viscosity?
betas=2.0
END SUBROUTINE SCENARIOS
=====
!
!
!
SUBROUTINE ADVANCE
!
! This subroutine advances the particles at one time-step
!
!
INCLUDE 'param.inc'
REAL vel0(nbody,dim)
REAL den0(nbody), hh0(nbody), u0(nbody)
! advance particles at first time-step
IF(tnow == 0.0)THEN
! make tree and find neighbors, smoothing
! length, and density
CALL TREENNSSL
! find all states of the system
CALL STATE
! find minimum time-step
CALL COURANT
DO i=1, nbody
IF(nnssl == 2) hh(i)=hh(i)+hhdot(i)*dtmin/2.0
IF(dsm == 2) den(i)=den(i)+dendot(i)*dtmin/2.0
IF(isorad == 1) u(i)=u(i)+udot(i)*dtmin/2.0
DO j=1, dim
vel(i,j)=vel(i,j)+acc(i,j)*dtmin/2.0
pos(i,j)=pos(i,j)+vel(i,j)*dtmin
END DO
END DO
tnow=tnow+dtmin
RETURN
ENDIF
! advance particles at first half time-step
DO i=1, nbody
hh0(i)=hh(i)
IF(nnssl == 2) hh(i)=hh(i)+hhdot(i)*dtmin/2.0
den0(i)=den(i)
IF(dsm == 2) den(i)=den(i)+dendot(i)*dtmin/2.0
u0(i)=u(i)
IF(isorad == 1) u(i)=u(i)+udot(i)*dtmin/2.0
DO j=1, dim
vel0(i,j)=vel(i,j)
vel(i,j)=vel(i,j)+acc(i,j)*dtmin/2.0
END DO
END DO
dtmin1=dtmin
! make tree and find neighbors, smoothing
! length, and density
CALL TREENNSSL
! find all states of the system
CALL STATE
! find minimum time-step
CALL COURANT
dtmin2=dtmin1/2.0+dtmin/2.0
! advance particles at second half time-step
DO i=1, nbody
IF(nnssl == 2) hh(i)=hh0(i)+hhdot(i)*dtmin2
IF(dsm == 2) den(i)=den0(i)+dendot(i)*dtmin2
IF(isorad == 1) u(i)=u0(i)+udot(i)*dtmin2
DO j=1, dim
vel(i,j)=vel0(i,j)+acc(i,j)*dtmin2
pos(i,j)=pos(i,j)+vel(i,j)*dtmin2
END DO
END DO

```



```

        ! link new cell into tree in place of old body
        subp(q,qind)=c
    ENDIF
    ! at this point, the node indexed by (q,qind) is known to
    ! be a cell, so advance to the next level of tree, and loop
    q=subp(q,qind)
    qind=SBINDX(p,q)
    GOTO 10
ENDIF
! found place in tree for p, so store it there
subp(q,qind)=p
END SUBROUTINE LDBODY
!=====
!////////////////////
!=====
    INTEGER FUNCTION MKCELL()
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!    This function allocates a cell and returns its index
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    ! check remaining space for a new cell
    IF(ncell > mxcell)THEN
        CALL ERROR('MKCELL: no more memory')
    ENDIF
    ! increment cell counter, initialize new cell pointer
    ncell=ncell+1
    MKCELL=ncell+nbody
    ! zero pointers to subcells of new cell
    DO i=1, nsubc
        subp(MKCELL,i)=0
    END DO
END FUNCTION MKCELL
!=====
!////////////////////
!=====
    INTEGER FUNCTION SBINDX(p,q)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!    This function computes subcell index for node p within cell q
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    INTEGER p, q
    ! initialize subindex to point to lower left subcell
    SBINDX=1
    ! loop over all spatial dimensions
    DO i=1, dim
        IF(pos(p,i) >= mid(q,i)) SBINDX=SBINDX+2**(dim-i)
    END DO
END FUNCTION SBINDX
!=====
!////////////////////
!=====
    SUBROUTINE TREEPROP
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!    This subroutine checks tree structure, assigns critical radius
!    for each cell, computes cell masses, c.m. positions, and
!    quadrupole moments
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    INTEGER p, q
    REAL pos0(dim), dist2
    ! list cells in order of descending size
    CALL SORTLIST
    ! loop processing cells from smallest to root
    DO i=ncell, 1, -1
        p=sortind(i)
        ! check that p is a cell
        IF(p < incell) CALL ERROR('TREEPROP: wrong cell')
        ! zero accumulators for this cell
        mass(p)=0.0
        pos0=0.0
        ! compute cell properties as sum of properties
        ! of its subcells
        DO j=1, nsubc
            q=subp(p,j)
            ! only access cells which exist
            IF (q /= 0)THEN
                ! sum properties of subcells to obtain
                ! values for cell p
                mass(p)=mass(p)+mass(q)
                DO k=1, dim

```



```

        pos0(k)=pos0(k)+mass(q)*pos(q,k)
    END DO
ENDIF
END DO
! normalize center of mass coordinates by total cell mass
DO j=1, dim
    pos0(j)=pos0(j)/mass(p)
END DO
! check tree, compute cm-to-mid distance
! and assign cell position
dist2=0.0
DO j=1, dim
    IF(pos0(j) < mid(p,j)-clsize(p)/2.0 .OR.
    &    pos0(j) >= mid(p,j)+clsize(p)/2.0)
    &    CALL ERROR('TREEPROP: tree structure error')
    dist2=dist2+(pos0(j)-mid(p,j))**2
    ! copy cm position to cell. This overwrites the midpoint
    pos(p,j)=pos0(j)
END DO
! assign critical radius for cell, adding offset
! from midpoint for more accurate forces. This
! overwrites the cell size
rcrit2(p)=(clsize(p)/theta+SQRT(dist2))**2
! compute quadrupole moments
DO j=1, nquad
    quad(p,j)=0.0
END DO
! loop over descendants of cell p
DO j=1, nsubc
    q=subp(p,j)
    IF(q /= 0)THEN
        ! sum properties of subcell q to
        ! obtain values for cell p
        DO m=1, MIN(2,dim)
            DO n=m, dim
                l=(m-1)*(dim-1)+n
                quad(p,l)=quad(p,l)+3.0*mass(q)*(pos(q,m)
    &                -pos(p,m))*(pos(q,n)-pos(p,n))
                IF(m == n)THEN
                    DO k=1, dim
    &                        quad(p,l)=quad(p,l)-mass(q)*(pos(q,k)
                        -pos(p,k))**2
                    END DO
                ENDIF
                ! if q itself is a cell, add its moments too
                IF(q >= incell) quad(p,l)=quad(p,l)+quad(q,l)
            END DO
        END DO
    END DO
END DO
END DO
END DO
END SUBROUTINE TREEPROP
=====
!////////////////////
!////////////////////
!////////////////////
=====
SUBROUTINE SORTLIST
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!    This subroutine sorts cells from largest (root) to smallest
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    INTEGER facell, lacell, nacell
    ! start scan with root as only active cell
    sortind(1)=root
    facell=1
    lacell=1
    ! loop while active cells to process
10  IF(facell <= lacell)THEN
        ! start counting active cells in next iteration
        nacell=lacell
        ! loop over subcells of each active cell
        DO i=1, nsubc
            DO j=facell, lacell
                ! add all cells on next level to active list
                IF(subp(sortind(j),i) >= incell)THEN
                    nacell=nacell+1
                    IF(nacell > maxn*nsubc)
    &                        CALL ERROR('SORTLIST: overflow')
                    sortind(nacell)=subp(sortind(j),i)
                ENDIF
            END DO
        END DO
    END IF

```

```

END DO
! advance first and last active cell indices, and loop
facell=lacell+1
lacell=nacell
GOTO 10
ENDIF
! above loop should list all cells; check the count
IF(nacell /= ncell)THEN
  WRITE(*,*) nacell, ncell
  CALL ERROR('SORTLIST: inconsistent cell count')
ENDIF
END SUBROUTINE SORTLIST
=====
!////////////////////
=====
SUBROUTINE NNSL
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
! This subroutine finds the nearest neighbors and smoothing
! length of particles
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
INCLUDE 'param.inc'
REAL DIST
INTEGER p, q, qsub
IF(nnssl == 1)THEN
  ! fixed smoothing length
  DO p=1, nbody
    hh(p)=1.5*(mass(p)/den(p))**(1.0/dim)
    ! effective radius according to hh
    IF(skf == 1) rp=3.0*hh(p)
    IF(skf == 2) rp=2.0*hh(p)
    IF(skf == 3) rp=3.0*hh(p)
    neighb(p)=0
    ! loop processing cells from root to smallest
    DO i=1, ncell
      q=sortind(i)
      rr=rp+clsize(q)
      IF(DIST(p,q,rr,0) < 0.0)THEN
        ! accepted: permit descent
        DO j=1, nsubc
          qsub=subp(q,j)
          IF(qsub /= 0 .AND. qsub < incell)THEN
            ! a body: skip self-consideration
            IF(qsub /= p)THEN
              ! test its spacing
              rr=rp
              IF(DIST(p,qsub,rr,1) < 0.0)THEN
                ! accepted as a nearest neighbor
                neighb(p)=neighb(p)+1
                ! check number of nearest neighbors
                IF(neighb(p) == nbbody)
                  & CALL ERROR('NNSL: too many')
                neighblist(p,neighb(p))=qsub
              ENDIF
            ENDIF
          ENDIF
        END DO
      END DO
      CALL SORTNEIGHB(p)
    END DO
  ELSEIF(nnssl == 2)THEN
    ! variable smoothing length
    DO p=1, nbody
      numiter=0
      numiter=numiter+1
      IF(numiter > 20)THEN
        WRITE(*,*) p, dennew, hhnew
        pause
        CALL ERROR('NNSL: too many iteration')
      ENDIF
      ! first use the smoothing length at this time, which
      ! is advanced via dh/dt=(h/dim)*divvel and find
      ! effective radius according to this hh
      IF(skf == 1) rp=3.0*hh(p)
      IF(skf == 2) rp=2.0*hh(p)
      IF(skf == 3) rp=3.0*hh(p)
      neighb(p)=0
      ! loop processing cells from root to smallest
      DO i=1, ncell
        q=sortind(i)

```

```

rr=rp+clsize(q)
IF(DIST(p,q,rr,0) < 0.0)THEN
  ! accepted: permit descent
  DO j=1, nsubc
    qsub=subp(q,j)
    IF(qsub /= 0 .AND. qsub < incell)THEN
      ! a body: skip self-consideration
      IF(qsub /= p)THEN
        ! test its spacing
        rr=rp
        IF(DIST(p,qsub,rr,1) < 0.0)THEN
          ! accepted as a nearest neighbor
          neighb(p)=neighb(p)+1
          ! check number of nearest neighbors
          IF(neighb(p) == nbody)
            CALL TERROR('NNSL: too many')
          neighblist(p,neighb(p))=qsub
        ENDIF
      ENDIF
    ENDIF
  END DO
END DO
END DO
! next find density by a summation over the particles
hmin=hh(p)
dennew=mass(p)*W(p,p)
DO jcursor=1, neighb(p)
  j=neighblist(p,jcursor)
  dennew=dennew+mass(j)*W(p,j)
END DO
! change the smoothing length via the
! proportionally (1/den)^(1/dim)
hhnew=2.0*(mass(p)/dennew)**(1.0/dim)
! check convergence of smoothing length
hfrac=ABS(hhnew-hh(p))/hh(p)
IF(hfrac > 0.01)THEN
  hh(p)=hhnew
  GOTO 10
ENDIF
CALL SORTNEIGHB(p)
END DO
ENDIF
END SUBROUTINE NNSL
=====
!////////////////////
!////////////////////
!////////////////////
FUNCTION DIST(i,q,rr,mode)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
! This function estimates the spacing criterion between particle
! p and node q
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
  INCLUDE 'param.inc'
  REAL DIST, rpq
  INTEGER q
  rpq=0
  IF(mode == 1)THEN
    DO j=1, dim
      a=pos(q,j)-pos(i,j)
      rpq=rpq+a*a
    END DO
    DIST=rpq-rr*rr
  ELSE
    DO j=1, dim
      a=ABS(pos(q,j)-pos(i,j))
      rpq=MAX(rpq,a)
    END DO
    DIST=rpq-rr
  ENDIF
END FUNCTION DIST
=====
!////////////////////
!////////////////////
!////////////////////
SUBROUTINE SORTNEIGHB(i)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
! This subroutine sorts the nearest neighbors at ascending
! distance to particle i
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
  INCLUDE 'param.inc'
  REAL rij(neighb(i))
  INTEGER indx(neighb(i)),indxn(neighb(i))

```

```

rij=0.0
DO jcursor=1, neighb(i)
  j=neighblist(i,jcursor)
  DO k=1, dim
    rij(jcursor)=rij(jcursor)+(pos(i,k)-pos(j,k))**2
  END DO
  rij(jcursor)=SQRT(rij(jcursor))
END DO
CALL INDEXX(neighb(i),rij,indx)
DO j=1, neighb(i)
  indx(j)=neighblist(i,indx(j))
END DO
DO j=1, neighb(i)
  neighblist(i,j)=indx(j)
END DO
END SUBROUTINE SORTNEIGHB

=====
!//////////////////////
!////////////////////////////////////////
=====
SUBROUTINE INDEXX(n,arr,indx)
!ccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
! This subroutine indexes an array arr(1:n), i.e. output the
! array indx(1:n) such that arr(indx(j)) is in ascending order
! for j=1,2,...,n. According to 'Numerical Recipes', Press et al.
!ccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
  INCLUDE 'param.inc'
  INTEGER n,indx(n),M,NSTACK
  REAL arr(n)
  PARAMETER (M=7,NSTACK=50)
  INTEGER i,indxt,ir,itemp,j,jstack,k,l,istack(NSTACK)
  REAL a
  DO j=1,n
    indx(j)=j
  END DO
  jstack=0
  l=1
  ir=n
1 IF(ir-l.lt.M)THEN
  DO j=l+1,ir
    indxt=indx(j)
    a=arr(indxt)
    DO i=j-1,l,-1
      IF(arr(indx(i)).le.a) GOTO 2
      indx(i+1)=indx(i)
    END DO
    i=l-1
    indx(i+1)=indxt
2 END DO
  IF(jstack.eq.0) RETURN
  ir=istack(jstack)
  l=istack(jstack-1)
  jstack=jstack-2
ELSE
  k=(l+ir)/2
  itemp=indx(k)
  indx(k)=indx(l+1)
  indx(l+1)=itemp
  IF(arr(indx(l)).gt.arr(indx(ir)))THEN
    itemp=indx(l)
    indx(l)=indx(ir)
    indx(ir)=itemp
  ENDIF
  IF(arr(indx(l+1)).gt.arr(indx(ir)))THEN
    itemp=indx(l+1)
    indx(l+1)=indx(ir)
    indx(ir)=itemp
  ENDIF
  IF(arr(indx(l)).gt.arr(indx(l+1)))THEN
    itemp=indx(l)
    indx(l)=indx(l+1)
    indx(l+1)=itemp
  ENDIF
  i=l+1
  j=ir
  indxt=indx(l+1)
  a=arr(indxt)
3 CONTINUE
  i=i+1
  IF(arr(indx(i)).lt.a) GOTO 3
4 CONTINUE

```



```

        vij(k)=vel(i,k)-vel(j,k)
        rij(k)=pos(i,k)-pos(j,k)
        rij0=rij0+rij(k)**2
        vijrij=vijrij+vij(k)*rij(k)
    END DO
    rij0=SQRT(rij0)
    IF(vijrij < 0.0)THEN
        muij=vijrij*hmin/(rij0**2+(eta*hmin)**2)
        phiij=(-alphas*vsig*muij+betas*muij**2)/denmin
    ELSE
        phiij=0.0
    ENDIF
    preden=pre(i)/den(i)**2+pre(j)/den(j)**2
    rdotdelW=0.0
    vdotdelW=0.0
    DO k=1, dim
        gradW=dW(i,j)*rij(k)/rij0
        acc(i,k)=acc(i,k)-mass(j)*(preden+phiij)*gradW
        rdotdelW=rdotdelW+rij(k)*gradW
        vdotdelW=vdotdelW+vij(k)*gradW
    END DO
    udot(i)=udot(i)+0.5*mass(j)*(preden+phiij)*vdotdelW
END DO
END SUBROUTINE RATES
!=====
!////////////////////
!=====
FUNCTION W(i,j)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!   This function evaluates the kernel of particles i and j
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    REAL sig, rij, q, W
    rij=0.0
    DO k=1, dim
        rij=rij+(pos(i,k)-pos(j,k))**2
    END DO
    rij=SQRT(rij)
    htest=(hh(i)+hh(j))/2.0
    IF(hmin /= hh(i) .AND. hmin /= htest)
&        CALL ERROR('KERNEL: hmin is inconsistent')
    q=rij/hmin
    ! Gauss kernel
    IF(skf == 1)THEN
        sig=(1.0/3.14)**(dim/2.0)
        IF(q <= 3.0)THEN
            W=EXP(-q*q)
        ELSE
            W=0.0
        ENDIF
        W=sig*W/hmin**dim
    ! spline-base kernel
    ELSEIF(skf == 2)THEN
        IF(dim ==1) sig=2.0/3.0
        IF(dim ==2) sig=10.0/(7.0*3.14)
        IF(dim ==3) sig=1.0/3.14
        IF(q <= 1.0)THEN
            W=1.0-1.5*q**2+0.75*q**3
        ELSEIF(q <= 2.0)THEN
            W=0.25*(2.0-q)**3
        ELSEIF(q > 2.0)THEN
            W=0.0
        ENDIF
        W=sig*W/hmin**dim
    ! Quintic kernel
    ELSEIF(skf == 3)THEN
        IF(dim ==1) sig=1.0/120.0
        IF(dim ==2) sig=7.0/(480.0*3.14)
        IF(dim ==3) sig=1.0/(120.0*3.14)
        IF(q <= 1.0)THEN
            W=(3.0-q)**5-6.0*(2.0-q)**5+15.0*(1.0-q)**5
        ELSEIF(q <= 2.0)THEN
            W=(3.0-q)**5-6.0*(2.0-q)**5
        ELSEIF(q <= 3.0)THEN
            W=(3.0-q)**5
        ELSEIF(q > 3.0)THEN
            W=0.0
        ENDIF
        W=sig*W/hmin**dim

```

```

ENDIF
END FUNCTION W
=====
!////////////////////
!
FUNCTION dw(i,j)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!   This function evaluates the differential of kernel
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    INCLUDE 'param.inc'
    REAL sig, rij, q, dw
    rij=0.0
    DO k=1, dim
        rij=rij+(pos(i,k)-pos(j,k))**2
    END DO
    rij=SQRT(rij)
    htest=(hh(i)+hh(j))/2.0
    IF(hmin /= hh(i) .AND. hmin /= htest)
&        CALL ERROR('KERNEL: hmin is inconsistent')
    q=rij/hmin
    ! Gauss kernel
    IF(skf == 1)THEN
        sig=(1.0/3.14)**(dim/2.0)
        IF(q <= 3.0)THEN
            dw=-(2.0*q*EXP(-q*q))/hmin
        ELSE
            dw=0.0
        ENDIF
        dw=sig*dw/hmin**dim
    ! spline-base kernel
    ELSEIF(skf == 2)THEN
        IF(dim ==1) sig=2.0/3.0
        IF(dim ==2) sig=10.0/(7.0*3.14)
        IF(dim ==3) sig=1.0/3.14
        IF(q <= 1.0)THEN
            dw=(-3.0*q+2.25*q**2)/hmin
        ELSEIF(q > 1.0 .AND. q <= 2.0)THEN
            dw=-0.75*(2.0-q)**2/hmin
        ELSEIF(q > 2.0)THEN
            dw=0.0
        ENDIF
        dw=sig*dw/hmin**dim
    ! Quintic kernel
    ELSEIF(skf == 3)THEN
        IF(dim ==1) sig=1.0/120.0
        IF(dim ==2) sig=7.0/(480.0*3.14)
        IF(dim ==3) sig=1.0/(120.0*3.14)
        IF(q <= 1.0)THEN
            dw=(-5.0*(3.0-q)**4+30.0*(2.0-q)**4-75.0*(1.0-q)**4)/hmin
        ELSEIF(q > 1.0 .AND. q <= 2.0)THEN
            dw=(-5.0*(3.0-q)**4+30.0*(2.0-q)**4)/hmin
        ELSEIF(q > 2.0 .AND. q <= 3.0)THEN
            W=(-5.0*(3.0-q)**4)/hmin
        ELSEIF(q > 3.0)THEN
            dw=0.0
        ENDIF
        dw=sig*dw/hmin**dim
    ENDIF
END FUNCTION dw
=====
!////////////////////
!
SUBROUTINE ERROR(message)
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
!   This subroutine stops the program if there is any error
!cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
    CHARACTER(*) message
    WRITE(*,*) '*****!?!?! error:!!?!*****'
    WRITE(*,*) message
    WRITE(*,*) '*****program is terminated*****'
    STOP
END SUBROUTINE ERROR
=====
!////////////////////
!
!=====
!   param.inc
!=====
!   this file contains common definitions and parameters
!=====

```



```

! dimension and maximum number of particles
INTEGER dim
! dim--> number of spatial dimensions (1, 2, or 3)
PARAMETER(dim=1)
INTEGER maxn
! maxn--> maximum number of SPH particles
PARAMETER(maxn=5000)
INTEGER nbody
! nbody--> number of SPH particles
REAL ul, ut, um
! ul--> unit of length (m)
! ut--> unit of time (s)
! um--> unit of mass (kg)
! ud--> unit of density (kg/m^3)
!   =um/ul^3
! uv--> unit of velocity (m/s)
!   =ul/ut
! ub--> unit of magnetic field (tesla)
!   =SQRT(um/ul)/ut
! G0--> gravitational constant
!   =(6.68e-11*um/ul)*(ut/ul)**2
COMMON /main/ nbody, ul, ut, um
! switches for different scenarios
INTEGER isorad
! isorad--> isothermal or adiabatic shock?
!   =1: adiabatic
!   =2: isothermal
INTEGER skf
! skf--> smoothing kernel function
!   =1: Gauss kernel (Gingold & Monaghan 1981)
!   =2: spline-base kernel (Monaghan 1985)
!   =3: Quintic kernel (Morris 1997)
INTEGER nnssl
! nnssl--> nearest neighbor search and smoothing length
!   =1: tree with h=hfac*(mass/den)**(1/dim)
!   =2: tree with dh/dt=(h/dim)*divvel
!   =3: tree with fixed neighbors between max and min
INTEGER dsm
! dsm--> density summation method
!   =1: summation model without continuity
!   =2: use continuity equation
COMMON /senarl/ isorad, skf, nnssl, dsm
! tolerance and correction parameters
REAL alphas, betas
! alphas--> shear viscosity
! betas--> bulk viscosity
REAL epsi, eta
! epsi--> parameter in XSPH correction of velocities
! eta--> parameter to avoid singularities in viscosity
PARAMETER(eps=0.5, eta=0.1)
REAL theta, eps
! theta--> tolerance parameter in tree structure
! eps--> tolerance parameter in tree structure
PARAMETER(theta=0.25, eps=1.0e-4)
REAL cour
! cour--> Courant number in step-time
PARAMETER(cour=0.25)
COMMON /toleran/ alphas, betas
! tree structure data arrays
INTEGER nsubc, nquad
! nsubc--> number of descendants per cell
! nquad--> number of independent quadrupole components
PARAMETER(nsubc=2**dim, nquad=2*dim-1)
INTEGER inode
! inode--> initial number of nodes (bodies + cells)
PARAMETER(inode=maxn+nsubc*maxn)
INTEGER mxcell, node, incell, ncell
! mxcell--> number of cells in the system (=nsubc*nbod)
! node--> number of nodes (bodies + cells)
! incell--> index of first cell in arrays (=nbod+1)
! ncell--> number of cells currently in use (<=mxcell)
INTEGER subp(inode,nsubc), root, sortind(maxn*nsubc)
! subp--> descendent of each cell
! root--> index of cell representing root (=incell)
! sortind(maxn*nsubc)--> sorted cells in descending size
REAL mid(inode,dim), clsize(inode)
! mid--> geometric center of each cell
! clsize--> size of each cell
REAL rcrit2(inode), quad(inode,nquad)
! rcrit2(incell:inode)--> critical distances^2 of each cell

```

```

! quad(incell:node,nquad)--> quad moments of each cell
COMMON /tree1/ mxcell, node, incell, ncell, sortind
COMMON /tree2/ rcrit2, quad, subp, root, mid, clsize, ndesc
! neighbor search parameters and smoothing length
INTEGER neighb(maxn), neighblist(maxn,maxn)
! neighb(maxn)--> number of neighbors for each particle
! neighblist(maxn,maxn)--> list of neighbors
REAL hh(maxn), hhdot(maxn)
! hh(maxn)--> smoothing lengths of SPH particles
! hhdot(maxn)--> smoothing length rate
REAL hmin
! hmin--> mean smoothing length of two neighbor particle
COMMON /neighbor1/ neighb, neighblist, hh, hhdot, hmin
! states of SPH particles
REAL pos(inode,dim), mass(inode), den(maxn)
! pos(node,dim)--> positions of bodies and cells

! mass(node)--> mass of bodies and cell
! den(maxn)--> density at position of each particle
REAL vel(maxn,dim), divvel(maxn)
! vel(maxn,dim)--> velocities of each body
! divvel(maxn)--> divergence of velocity

REAL acc(maxn,dim), dendot(maxn)
! acc(maxn,dim)--> acceleration of bodies
! dendot(maxn)--> density rate
REAL sound(maxn), pre(maxn), temp(maxn)
! sound(maxn)--> sound speed of particles
! pre(maxn)--> pressure of particles
! temp(maxn)--> temperature of particles
REAL u(maxn), udot(maxn)
! u(maxn)--> energy of particles
! udot(maxn)--> energy rate
COMMON /state1/ pos, mass, vel, divvel, acc, dendot, u, udot
COMMON /state2/ den, sound, pre, temp
! time integration parameters
REAL tnow, dtmin
! tnow--> current time
! dtmin--> minimum time-step
COMMON /time/ tnow, dtmin
=====
!=====

```